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GenCore version 5.1.6 copyright (c) 1993 - 2003 Compugen Ltd.
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- nucleic search, using frame\_plus\_p2n model

OM protein

August 29, 2003, 09:51:49; Search time 2055 Seconds (without alignments) 1596.644 Million cell updates/sec Run on:

US-09-866-582A-2 661 1 MDAVDRGVYFDEDFHVRILD......GEQELMIQKLSDSSSGAAYV 135 Perfect score;

0.5 7.0 7.0 Xgapop 10.0, Xgapext (Ygapop 10.0, Ygapext (Fgapop 6.0, Fgapext Delop 6.0, Delext **BLOSUM62** Scoring table: Sequence:

22781392 segs, 12152238056 residues Searched: Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0% .Maximum Match 100% Listing first 100 summaries

Command line parameters:
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-FGAPEXT=7 -YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7

Database :

em\_gss\_fun:\* em\_gss\_vrt:, \_gss\_mam: em\_gss\_hum: \_gss\_inv: em\_gss\_pln: em\_gss\_pro: em\_estba:\*
em\_estum:\*
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em\_gss\_vrl:\* gb\_gssl:\*

em\_gss\_phg:

em\_gss\_rod:

29: gb\_gss2:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

scription		837 AV3928	2290 894059C	0429 BW28042	0674 BW2806	6060 BP02606	246 AV964246	5694 BW2156	3658 BP01365	18971 BP00897	18/82 BWZ98/8	3621 BW31362	9944 BWU/994	0900 BW10090	2150 BW01/43	763 AVR97763	2507 BW3125	3735 NISC_nq	.937 AL79193	467 AL64946	277 AL637277	15126 BY63512	0890 BWIDU8	426 AV899426	8613 BW24861	16639 BW1	7926 BW11792	4012 BJ06401	1392 CM34IIU.	1692 UI-R-ABI-	6900 ug64h07	18168 BY43816	1561 UI-R-E	19218 UI-K-DK	584 ui88c05.	881	6762 ih70c01	38726 UI-R-DY	4/85 UI-M-DD	1938 AGENC	7210 AGENC	)9888 UI-R-	70290 H3085G04	1092/ UI-M-CG	/222 UI-M-DUZ	3210 H3085G0	2397 Mus mus	56846 ut88d02.	36260 uf76e04	6815 ih71c0	4/31 BAZ/4/31	1517 pq11n.p	7180 60417225	9785 BX25978	AV882930 AV882930 BU409809 603159073
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BX260830 BX260830 BU490085 604129871 BU11527 603138847 BU346945 604170279 BU3414994 603667387 BU357913 603407539 BU357913 603407539 BU35505 603412313 BU315209 603165294 BU108921 60316316 BU108921 603103889 BU1077813 603473673 BU1077813 603113889 BU37748 603535963	BU397453 603535302 BU458001 603777109 BG101052 UY18908.Y BU305614 603612226 BB789469 BB789469 BF075390 224382 MA AA059304 Z4878605.r AA273622 va23204.r CA941070 ir38b02.y BF075374 224358 MA B171111 ie07h04.y BC632674 i126d03.y BX096677 BX099677 BX413118 BU914040 AGENCOURT N40217 yx993803.r1 BU074070 in28c01.y BJ5029640 BJ029640	2 2 2 2	Chlorophyta; Chlorophyceae; Volvocales; Chlorophyta; Chlorophyceae; Volvocales; Cato,S., Fukuzawa,H. and Tabata,S. nalysis of CDNAs in a unicellular green ridtli. I. Generation of 3433	esearch , Japan ://www.kazusa.or.jp/en/plant/. nhardtii"
29.3 742 11.229.3 7.24 12.29 12.2	194 29.3 1808 13 B0397453 193 29.2 422 13 B0301052 19.2 29.2 422 10 BG101052 19.2 29.1 428 10 BF078364 19.2 29.0 391 10 BF075390 19.2 29.0 446 9 AAV597208 19.2 29.0 457 9 AAV597304 19.2 29.0 457 9 AAV597304 19.2 29.0 653 10 BF075374 19.2 29.0 611 12 B1713151 19.2 29.0 626 13 BX099677 19.2 29.0 876 13 BX099677 19.2 29.0 876 13 BX099677 19.3 28.9 510 13 BX0974040	191 28.9 531 10 BF075368 191 28.9 647 14 CA381325 191 28.9 649 13 BX419276 191 28.9 1050 13 BX419276 ALIGNMENTS AV392837 . 466 bp mF	AV392837 Chlamydomonas reinhardtii C9 Chlamydomonas CDNA clone CM100a09_r 5', mRNA sequence. AV392837.1 GI:6547053 EST. Chlamydomonas reinhardtii Chlamydomonas reinhardtii Chlamydomonas reinhardtii Eukaryota; Viridiplantae; Chlorophyta; Chlorophycea Chlamydomonadaceae; Chlamydomonas. 1 (bases 1 to 466) Asamizu.E., Nakamura,Y., Sato,S., Fukuzawa,H. and TA large scale structural analysis of cDNAs in a uni alga, Chlamydomonas reinhardtii. I. Generation of 3 non-redundant expressed sequence tags	July288 Contact: Yasukazu Nakamura The First Laboratory for Plant Gene Research Takazusa DNA Research Institute Yana 1532-3, Kisarazu, Chiba 292-0812, Japan Emall: Ynakamu@kazusa.or.jp, URL:http://www.k. Location/Qualifiers 1466 /organism="Chlamydomonas reinhardtii //mol_type="nRNA" //strain="c9" //db_xref="taxon:3055"

RESULT 1 AV392837 LOCUS DEFINITION

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ACCESSION VERSION KEYWORDS SOURCE ORGANISM

REFERENCE AUTHORS TITLE JOURNAL MEDLINE PUBMED COMMENT FEATURES SOUICE  $\sim$ 

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BW280674 Satch unpublished cDNA library, gonad Ciona intestinalis cDNA clone cigd014j07 5', mRNA sequence.
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                                                                                                                                                                                                                                                                                           7 GlyValTyrPheAspGluAspPheHisValArgIleLeuAspValAspLysTyrAsnAla 26
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Tel: 81-75-753-4081
Fax: 81-75-705-1113
Email: satoh@ascidian.zool.kyoto-u.ac.jp.
Location/Qualifiers
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Department of Zoology
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                                                                                                                    Anote="Vector: pBluescript II SK-; Site_1: EcoRI; Site_2:
XhOI; This library, constructed by John Davies and Jeffrey
McDermott, combines cDNAs from CC-1690 cells grown to
mid-log phase in TAPP (acetate-containing) medium in the
light, TAP medium in the dark, HS (minimal) medium in the
light, TAP medium in the dark, HS (minimal) medium in
ambient levels of CO2 and HS medium bubbled with 5% CO2.
POlyA mRNA was purified from each sample, pooled and cDNA
synthesized. The cDNA was directionally cloned into lambda
ZAP II (Stratagene) in the EcoRN (5') and XhoI (3') sites.
pBluescript II SK- plasmids were excised from the lambda
ZAP clones by superinfection with ExAssist (Stratagene)
phage. The library was normalized using method 4 described
in Bonaldo et al (1996) Genome Research 6: 791-806."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  646 bp mRNA linear EST 11-NOV-2002 BW280429 Nori Satoh unpublished cDNA library, gonad Ciona intestinalis cDNA clone cigd013n23 5', mRNA sequence. BW280429 GI:24861040 EST.
                                                                              /clone_lib="C. reinhardtii CC-1690, normalized, Lambda Zap
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Bukaryota: Metazoa; Chordata; Urochordata; Ascidiacea; Enterogona;
Phlebobranchia: Cionidae; Ciona.
1 (bases 1 to 646)
Satou.X., Shin-i.T., Kohara,Y. and Satoh,N.
Expressed genes in Ciona intestinalis (2002c)
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Sakyo-ku, Kyoto, Kyoto 606-8502, Japan
Tel: 81-75-753-4081
Fax: 81-75-705-1113
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Department of Zoology
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1 (bases 1 to 540)
Inaba,K., Padma,P., Satouh,Y., Shin-i,T., Kohara,Y., Satoh,N. and
                                                                                                                                                                                                                                                                                                                                                LeuLysalaIleGlyLeuArgAsnArgValAlaAlaLeuSerGluGluArgLysArgLys 86
/tissue_type="gonad"
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1 108 c 121 g 196 t 3 others
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EST analysis of gene expression in testis of the ascidian Ciona intestinalis
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/db_xref="taxon:7719"
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Contact: Kazuo Inaba
Asamushi Marine Biological Statio
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Asamushi, Aomori, Aomori 039-3501,
Tel: 81-17-752-3394
Fax: 81-17-752-2765
                                                                                                                               Indels:
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14-MAR-2002 ::: |||::::::: ||||
143 ACATCTGAACTGAAAGAATGCCACGAATTCGTCTCGAAAATTCGACCAGTTTCAAAAA 202 Eukaryota; Metazoa; Chordata; Urochordata; Ascidiacea; Enterogona; Phlebobranchia; Cionidae; Ciona. I chases I to 541)
Satoh, N., Satou, Y., Kohara, Y. and Shin-i, T.
Expressed genes in Ciona intestinalis 7 GlyValTyrPheAspGluAspPheHisValArgIleLeuAspValAspLysTyrAsnAla 26 SerLysSerLeuGlnAspAsnThrAsnValPheIleAsnAsnIleGlnAsnMetGlnGly 46 /tissue\_type="testis" /clone\_lib="K. Inaba unpublished cDNA library, testis" 101 c 102 g 147 t 1 others 67 LeuLysAlaIleGlyLeuArgAsnArgValAlaAlaLeuSerGluGluArgLysArgLys LeuValAspLysTyrValSerAlaIleAspGlnGlnValGluArgLeuGluAlaGluLys AV964246 Nori Satoh unpublished cDNA library, egg Ciona intestinalis cDNA clone cieg17g02 5', mRNA sequence. AV964246 540 41 34 47 0 Sakyov (u. Kyoto, Kyoto 606-8502, Japan Sakyov (u. Kyoto, Kyoto 606-8502, Japan Tel: 81-75-753-4081 Fax: 81-75-705-1113 Email: satch@ascidian.zool.kyoto-u.ac.jp. Location/Qualifiers Conservative: /organism="Ciona intestinalis" /mol\_type="msNA" /db\_xref="taxon:7719" /clone="cieg17902" Mismatches: Indels: Length: Matches: US-09-866-582A-2 (1-135) x BP026060 (1-540) AV964246.1 GI:19453942 Unpublished Contact: Nori Satoh Department of Zoology 4.01e-08 200.00 61.48% 33.61% 30.26% Ciona intestinalis Ciona intestinalis Kyoto University 127 AspSer 128 443 ACTTCT 448 Percent Similarity: Best Local Similarity: σ 189

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Eukaryota; Metazoa; Chordata; Urochordata; Ascidiacea; Enterogona;
Phlebobranchia; Cionidae; Ciona.
1 (bases 1 to 565)
Satou,Y., Shin-1,T., Kohara,Y. and Satoh,N.
Expressed genes in Ciona intestinalis (2002c)
Unpublished
Contact: Nori Satoh
Department of Zoology
Kyoto University
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97 c 103 g 153 t
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Mismatches:
Indels:
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Location/Qualifiers
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Tel: 81-75-753-4081
Fax: 81-75-705-1113
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BP013658 Nori Satch unpublished cDNA library, young adult Ciona intestinalis cDNA clone ciad57n03 5', mRNA sequence.
BP013658.1 GI:19505135
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Satoh, N., Satou, Y., Kohara, Y. and Shin-i, T.
Expressed genes in Ciona Intestinalis
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/tissue_type="whole animal"
/dev_stage="egg"
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Location/Qualifiers
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Contact: Nori Satoh
Department of Zoology
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1 (bases 1 to 670)

2atch, N., Satou, Y., Kohara, Y. and Shin-i, T.

Expressed genes in Ciona intestinalis
Unpublished
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Indels:
Gaps:
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Sakyo-ku, Kyoto, Kyoto 606-8502, Japan
19-18-17-57-57-34081
Fax: 81-75-705-1113
Email: satoh@ascidian.zool.kyoto-u.ac.jp.
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Department of Zoology
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BW298785
BW298785 Nori Satoh unpublished cDNA library, neural complex Ciona intestinalis cDNA clone cinc003m02 5', mRNA sequence.
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I (bases I to 600)
Satou, Y., Shin i, T., Kohara, Y. and Satoh, N.
Expressed genes in Ciona intestinalis (2002c)
Contact: Nori Satoh
Department of Zoology
Kyoto University
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Conservative:
Mismatches:
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Location/Qualifiers
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Tel: 81-75-753-4081
Fax: 81-75-705-1113
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/clone="ciad50e21"
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BW298785.1 GI:24879396
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Best Local Similarity:
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Alignment Scores:
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Ciona intestinalis
Eukaryota; Metazoa; Chordata; Urochordata; Ascidiacea; Enterogona;
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1 (bases 1 to 683)
Satou, Y., Shin-i,T., Kohara,Y. and Satoh,N.
Unpublished
                                                                                                                 Conservative:
Mismatches:
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Location/Qualifiers
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Tel: 81-75-753-4081
Fax: 81-75-705-1113
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/db_xref="taxon:7719"
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Department of Zoology
Kyoto University
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BW079944 Nori Satoh unpublished cDNA linear EST 20-OCT-2002 intestinalis cDNA clone rcieg080kl3 3', mRNA sequence. BW079944 BW079944.1 GI:24181356
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Ciona intestinalis
Bukaryota; Metazoa; Chordata; Urochordata; Ascidiacea; Enterogona;
Phlebobranchia; Cionidae; Ciona
1 (bases 1 to 715)
Satou,Y., Shin-i,T., Kohara,Y. and Satoh,N.
Expressed genes in Ciona intestinalis (2002c)
                                                                                                                                                                                                                                                                                                                                                                                  63 GGGCTTCATTTTGACGAGTTAACAAGCTCAGAGTTTTAAAAACCAGACACTAATCAGCAG 122
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/db_xref="taxon:7719"
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108 c 123 g 210 t
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Sakyo-ku, Kyoto, Kyoto 606-8502, Japan
Bell 81-75-753-4081
Fax: 81-75-705-1113
Email: satch@ascidian.zool.kyoto-u.ac.jp.
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/clone="rcinc003m02"
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1 (basea: 1 to 719)
Satou, Y., Shin-i, T., Kohara, Y., and Satoh, N.
Expressed genes in Ciona intestinalis (2002c)
Unpublished
Contact: Nori Satoh
Department of Zoology
Kyoto University
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Conservative:
Mismatches:
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Tel: 81-75-753-4081
Fax: 81-75-705-1113
Email: satch@ascidian.zool.kyoto-u.ac.jp.
Location/Qualifiers
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/mol_type="mRNA"
/db_xref="taxon:7719"
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BW077451 Nori Satch unpublished cDNA library, egg Clona intestinalis cDNA clone rcieg072121 3', mRNA sequence. BW077451 GI:24178863
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bukaryota; Metazoa; Chordata; Urochordata; Ascidiacea; Enterogona;
Phlebobranchia; Cionidae; Ciona
1 (bases 1 to 720)
Sadou, Y., Shin-i, T., Kohara, Y. and Satoh, N.
Expressed genes in Ciona intestinalis (2002c)
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548 CTTGTTGGTAGCTTCATAGAGATGACCGACGCAATCGCAAAAGATGTTGAAAATGAGAAA 489
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/tissue_type="neural complex"
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complex" 259 t
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Sakyo-ku, Kyoto, Kyoto 606-8502, Japan
Tel: 81-75-753-4081
Fax: 81-75-705-1113
Email: Satoh@ascidian.zool.kyoto-u.ac.jp.
Location/Qualifiers
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                                                                                                                                                           Length:
Matches:
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Department of Zoology
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Department of Zoology
Kyoto University
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1 (bases 1 to 'Satou'Y.' Shin-17.7, Kohara,Y. and Satoh,N.
Expressed genes in Ciona intestinalis (2002c)
Unpublished
Contact: Nori Satoh
Department of Zoology
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/tissue_type="whole animal"
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Matches:.
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Location/Qualifiers
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/db_xref="taxon:7719"
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Tel: 81-75-753-4081
Fax: 81-75-705-1113
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BW213159
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AV897763 Nori Satoh unpublished cDNA library, young adult Ciona intestinalis cDNA clone rciad50e21 3', mRNA sequence.
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Eukaryota; Metazoa; Chordata; Urochordata; Ascidiacea; Enterogona;
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1 (bases 1 to 758)
Satob, N., Satou, Y., Kohara, Y. and Shin-i, T.
Expressed genes in Ciona intestinalis
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Tel: 81-75-753-4081
Fax: 81-75-705-1113
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us-09-866-582a-2.rst

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Eukaryota; Metazoa; Chordata; Urochordata; Ascidiacea; Enterogona;

Eukaryota; Metazoa; Chordata; Urochordata; Ascidiacea; Enterogona;

1 (bases 1 to 759)

Satou,Y., Shin-i,T., Kohara,Y. and Satoh,N.

Expressed genes in Ciona intestinalis (2002c)

Unpublished

Contact: Nori Satoh

Department of Zoology

Myoto University

Contact: Nori Satoh

Department of Loology

Myoto University

Contact.
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619 ACATCTGAAACTGAAAATGCCACGAATTCGTCTCGAAAAATCGACCAGTTTCAAAAA
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Location/Qualifiers
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Tel: 81-75-753-4081
Fax: 81-75-705-1113
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/mol_type="mRNA"
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Silurana tropicalis cDNA clone
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Silurana tropicalis
Silurana tropicalis
Silurana tropicalis
Silurana tropicalis
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea; Pipidae;
Xenopodinae; Silurana
1 (bases 1 to 615)
NIH-XCG http://image.llnl.gov/image/html/xenopuslib_info.shtml.
National institute of Child Health and Human Development, National
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ::: |||||||||| ||| ::: ||||||| 354 CAACACCAGTCTAAGAAACAAGAGGCCTGAACATGAGTTGAATTGAATTGAATTGAATTGAATTGTCA 413
                                                                                                                                                                                                                                                                                                                                                                                                                                 114 ACATCTGAACTGAAAGAGGAATGCCACGAATTCGTCTCGAAAATCGACCAGTTTCAAAAA 173
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                                                unpublished cDNA library, heart"
238 t
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cDNA Library Preparation:
CDNA Library Arrayed by: The I.M.A.G.E. Consortium/LLNL
DNA Sequencing by: National Institutes of Health Intramural
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41
34
47
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                                                                                                                                                                                    Length:
Matches:
Conservative:
Mismatches:
Indels:
/db_xref="taxon:7719"
/clone="ciht023ell"
/tissue_type="heart"
/clone_lib="Nori Satch un
a 113 c 133 g 238
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Contact: Robert Strausberg, Ph.D.
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IMAGE:5382476 5', mRNA sequence.
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Best Local Similarity:
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AL649467 XGC-gastrula Silurana tropicalis cDNA clone TGas049d15 5',
                                                                                                                                                                                                                                                                            /dev_stage="neurula"
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/clone_lib="XGCneurula"
/note="Vector: pC5107; Site_1: EcoRI; Site_2: NotI; cDNA
was oligo dT primed from 5ug of poly A+ RNA from neurula.
EcoRI-NotI cut cDNA was then ligated into pCS107 with
ECORI at the 5' end and NotI at the 3' end."
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                                                        Email: trop@sanger.ac.uk
Sanger Xenopous tropicalis EST project 2001
TROPICALIS_SEQUENCE_ID: TNeu094a07.plcSP6
Sequencing primer: PlCSP6
This sequence is from a Xenopus Gene Collection (XGC) library
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      619
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Matches:
Conservative:
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Silurana tropicalis
                                                                                                                                                                                                       /organism="Silurana tropicalis"
/mol_type="mRNA"
/db_xref="taxon:8364"
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                                      Hinxton, Cambridgeshire, CB10 1SA, UK
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                                                                                                                                                 constructed by Aaron M. Zorn.
Location/Qualifiers
1. .619
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AL649467
AL649467.1 GI:17658653
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   Contact: Taylor R
                       Sanger Centre
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AL649467
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                                                                                                                         /tissue_type="neurula"
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Silurana tropicalis
Subraryota: Metacoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia: Batrachia; Anura; Mesobatrachia; Pipoidea; Pipidae;
Xenopodinae; Silurana.
1 (bases 1 to 619)
1 Taylor, R., Ashurst, J.L., Croning, M.D.R., Zorn, A.M. and Rogers, J. Sanger Xenopus tropicalis EST project 2002
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Matches:
Conservative:
Mismatches:
Indels:
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/mol_type="mknA"
/db_xref="taxon:8364"
/clone="IMAGE:5382476"
Seq primer: M13RP1 reverse primer (ABI).
Location/Qualifiers
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291 GCCCAGCAGCAACTGTATGCCTTAATAGCAGAGAAAAATGCAACTAGAAAGGTAC 350
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /clone_lib="XGC-neurula"
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/note="Vector: pCS107; Site_1: EcoRI; Site_2: NotI; cDNA
was oligo dT primed from 5ug of poly A+ RNA from neurula.
EcoRI-NotI cut cDNA was then ligated into pCS107 with
EcoRI at the 5' end and NotI at the 3' end."
135 c 143 g. 155 t
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Silurana tropicalis
Silurana tropicalis
Subraryotas (horazos, chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachla; Anura; Mesobatrachla; Pipoidea; Pipidae:
Xenopodinae; Silurana.
I (bases I to 640)
Huckle, E., Taylor, R., Ashurst, J.L., Zorn, A.M. and Rogers, J.
Sanger Xenopus tropicalis EST project 2001 (10_2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hinxton, Cambridgeshire, CB10 1SA, UK
Email: trop@sanger.ac.uk
Sanger Xenopus tropicalis EST project 2001
TROPICALIS_ESQUENCE_ID: TNeu006k05.sp6
Sequencing primer: SP6
This sequence is from a Xenopus Gene Collection (XCC) library
constructed by Aaron M. Zorn.
Location/Qualifiers
1. 640
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/mol_type="mkna" silurana tropicalis"
/mol_type="mkna" silurana tropicalis"
/mol_type="mkna" silurana tropicalis"
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/dev_stage="neurula"
/lab_host="Escherichia coli DH10B"
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// Organism="Silurana tropicalis"
// Organism="RNA"
// Ob_xref="taxon:8364"
// Clone="TGas049415"
// Clone="TGas049415"
// Iab_host="gastrula (stages 10.5-13 mixed)"
// Iab_host="Escherichia coli XL1-blue"
// Clone="lb="XGC-gastrula"
// Iab_host="Escherichia coli XL1-blue"
// Clone="Vector: pCS107" site_1: EcoRI; Site_2: NotI: cDNA / note="Vector: pCS107" site_1 in Coli X Ar RNA from stages was oligo dT primed from 5ug of poly A+ RNA from stages 10-13 gastrulae. EcoRI-NotI cut cDNA was then ligated into pCS107 with EcoRI at the 5' end and NotI at the 3'
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1 (bases 1 to 629)
Huckle, E., Taylor, R., Ashurst, J. L., Zorn, A. M. and Rogers, J. Sanger Xenopus tropicalis EST project 2001 (10_2001)
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                                                                                                                                                                                                                                                                                                         Email: trop@sanger.ac.uk
Sanger Xenopus tropicalis EST project 2001
Sanger Xenopus tropicalis EST project 2001
Sanger Xenopus Tropicalis EST project 2001
Sequencing primer: SP6
This sequence is from a Xenopus Gene Collection (XGC) library constructed by Aeron M. Zorn.
Location/Qualifiers
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Division of Experimental Animal Research in Riken contributed to prepare mouse tissues.

Tissues were provided by Michela Fagiolini and Takao K. Hensch (Laboratory for Neuronal Circuit Development Brain Science Institute RIKEN 2-1 Hirosawa, Wako-shi, Saitama 351-0198 Japan ) whose assistance we gratefully acknowledge.

Please visit our web site (http://genome.gsc.riken.go.jp) for
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encyclopedia: real-time sequence clustering for construction of a nonredundant cDNA library. Genome Res. 11 (2), 281-289 (2001) CDNA library was prepared and sequenced in Mouse Genome Encyclopedia Project of Genome Exploration Research Group in Riken Genomic Sciences Center and Genome Science Laboratory in RIKEN.
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/tissue_type="visual cortex"
/clone_lib="RIKEN full-length enriched, visual cortex"
70 c 120 g 94 t
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/mol_type="mRNA"
/strain="C57BL/6J"
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Ciona intestinalis
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URL:http://genome.gsc.riken.go.jp/
Alzawa,K., Akimura,T., Arakawa,T., Carninci,P., Fukuda,S., Hirozane
Alzawa,K., Akimura,T., Arakawa,T., Konno,H., Miyazaki,A.
T., Imotani,K., Ishii,Y., Itoh,M., Kawai,J., Konno,H., Miyazaki,A.,
Murata,M., Nakamura,M., Nomura,K., Numazaki,R., Ohno,M., Sakai,K.,
Sakazume,N., Sasaki,D., Sato,K., Shibata,K., Shiraki,T., Tagami,
M., Waki,K., Watahiki,A., Muramatsu,M. and Hayashizaki,Y. Direct
Submission
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Nature 420, 563-573 (2002)
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The Institute of Physical and Chemical Research (RIKEN)
1-7-22 Suehiro-cho, Tsurumi-ku, Yokohama, Kanagawa 230-0045, Japan Tel: 81-45-503-9216
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              prepare full-length cDNA libraries for rapid discovery of new genes. Genome Res. 10 (10), 1617-1630 (2000)
KIKBN integrated sequence analysis (RISA) system--384-format sequencing pipeline with 384 multicapillary sequencer. Genome Res. 10 (11), 1757-1771 (2000)
Computer-based methods for the mouse full-length cDNA
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Mammalla; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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BY635126 RIKEN full-length enriched, visual cortex Mus musculus
cDNA clone K430066M20 3', mRNA sequence.
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Normalization and subtraction of cap-trapper-selected cDNAs
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411 TTTAAC 416
                                  125 LeuSer 126
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TITLE

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84

us-09-866-582a-2.rst

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Alignment Scores:
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BW286181.1 GI:24866792
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Ciona intestinalis
Eukaryota; Metazoa; Chordata; Urochordata; Ascidiacea; Enterogona;
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Eukaryota; Metazoa; Chordata; Urochordata; Ascidiacea; Enterogona;
                                                                                                                                                                                                    /organism-"Ciona intestinalis"
/mol_type="mRNA"
/db_xref="*axon.7719"
/clone="rcigd013n23"
/tissue_type="gonad"
/clone_lib="Nori Satoh unpublished cDNA library, gonad"
125 c 108 g 255 t 3 others
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    6 ArgGlyValTyrPheAspGluAspPheHisValArgIleLeuAspValAspLysTyrAsn
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                                                                                                                                                                                                                                                                                                                                           715
41
34
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0
            Phlebobranchia; Cionidae; Ciona.

1 (Dassa 1 to 715)

1 (Dassa 1 to 715)

Expressed genes in Ciona intestinalis (2002c)
Unpublished
Contact: Nori Satch
Department of Zoology
                                                                                                                                                                                                                                                                                                                                          Length:
Matches:
Conservative:
Mismatches:
Indels:
                                                                                                          Kyoto University
Sakyo-ku, Kyoto, Kyoto 606-8502, Japan
Tel: 81-75-753-4081
Fax: 81-75-705-1113
Fmail: satch@ascidian.zool.kyoto-u.ac.jp.
Location/Qualifiers
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Best Local Similarity:
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Ciona intestinalis
Eukaryota; Metazoa; Chordata; Urochordata; Ascidiacea; Enterogona;
Phlebobranchia; Cionidae; Ciona.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               7 GlyValTyrPheAspGluAspPheHisValArgIleLeuAspValAspLysTyrAsnAla 26
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/db_xref="taxon:7719"
/clone="cigdOgonad"
/tissue_type="gonad"
/clone_lib="Nori Satoh unpublished cDNA library, gonad"
a 104 c 116 g 191 t
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Phlebobranchia; Cionidae; Ciona.
1 (bases 1 to 633)
Satou,Y., Shin-i,T., Kohara,Y. and Satoh,N.
Uppublished
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40
47
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative:
Mismatches:
Indels:
                                                                                                                                                                                                                  Email: satoh@ascidian.zool.kyoto-u.ac.jp.
Location/Qualifiers
                                                                                                                                                       Sakyo-ku, Kyoto, Kyoto 606-8502, Japan
Tel: 81-75-753-4081
Fax: 81-75-705-1113
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length:
Matches:
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AV899426.1 GI:16888524
                                                                                                   Contact: Nori Satoh
Department of Zoology
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1.01e-07
196.00
61.48%
32.79%
29.65%
                                                                                                                                           Kyoto University
                                                                                                                                                                                                                                                             1. 633
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Best Local Similarity:
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238
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BW156639/c
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BW248613.1 GI:24828531
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Ciona intestinalis
Eukaryota; Metazoa; Chordata; Urochordata; Ascidiacea; Enterogona;
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                                                                                                                            Email: satch(dasidian.zool.kyoto-u.ac.jp.
Location/qualifiers
1. 661
/organism="ciona intestinalis"
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/clone="rciad74dm19"
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/dev_stage="young adult"
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                                                                                                                                                                                                                                                                                                                                                                                                                                          8 ValTyrPheAspGluAspPheHisValArgIleLeuAspValAspLysTyrAsnAlaSer
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                                                                                                                                                                                                                                                                           3 others
1. (bases 1 to 661)
Satoh, N., Satou, Y., Kohara, Y. and Shin-i, T.
Expressed genes in Ciona intestinalis
Unpublished
Contact: Nori Satoh
Department of Zoology
Kyoto University
Sakyo-ku, Kyoto, Kyoto 606-8502, Japan
Tel: 81-75-753-4081
Fax: 81-75-705-1113
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Matches:
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196.00
61.16%
33.06%
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TCT 297
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BW248613
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BW156639 Nori Satch unpublished cDNA library, gonad Ciona intestinalis cDNA clone rcigd002d11 3', mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                             /tissue_type="whole animal"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   65 GGGCTTCATTTTGACGAAGTTAACAAGCTCAGAGTCTTAAAACCAGAGACTAATCAGCAG 124
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Phlebobranchia, Cionidae, Ciona.

1 (bases 1 to 674)
Satou,Y., Shin-i,T., Kohara,Y. and Satoh,N.
Expressed genes in Ciona intestinalis (2002c)
Contact: Nori Satoh
Department of Zoology
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Matches:
Conservative:
Mismatches:
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Location/Qualifiers
1.674 /organism="Ciona intestinalis"
                                                                                                                                             Kyoto University
Sakyo-ku, Kyoto, Kyoto 606-8502, Japan
Tel: 81-75-753-4081
Fax: 81-75-705-1113
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BW156639.1 GI:24513864
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Query Match:
DB:
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Phlebobranchia; Cionidae; Ciona
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BW117926 GI:24364591
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Ciona intestinalis
Eukaryota; Metazoa; Chordata; Urochordata; Ascidiacea; Enterogona;
Eukaryota; Metazoa; Chordata; Urochordata; Ascidiacea; Enterogona; Phlebobranchia; Cionidae; Ciona.

1 (bases 1 to 718)

Satou, Y., Shin-i, T., Kohara, Y. and Satoh, N.

Expressed genes in Ciona intestinalis (2002c)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     130 CAACAACAGTTGAAAGCACTCATAGCGGAGAAGAAAAACACAACTTGAGCGATTGAATGTG 371
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  550 CTTGTTGGTAGCTTCATAGAAATGACCGACGCAATCGCAAAAGATGTTGAAAATGAGAAA 491
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/clone_lib="Nori Satoh unpublished cDNA library, gonad"
127 c 110 g 260 t
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Matches:
Conservative:
Mismatches:
                                                                                                                           Sakyo-ku, Kyoto, Kyoto 606-8502, Japan
Tel: 81-75-753-4081
Fax: 81-75-705-1113
Email: satch@ascidian.zool.kyoto-u.ac.jp.
Location/Qualifiers
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/organism="Ciona intestinalis"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    (1-718)
                                                                                                                                                                                                                           /mol_type="mRNA"
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/clone="rcigd002d11"
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                                                                                    Contact: Nori Satoh
Department of Zoology
Kyoto University
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BW117926/c
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AUTHORS
TITLE
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BJ064012
BJ064012 NIBB Mochii normalized Xenopus tailbud library Xenopus laevis cDNA clone XL078b07 5', mRNA sequence.
BJ064012
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                                                                                                                                                                                                                                                                                                                                                     /tissue_type="whole animal"
/dev_stage="tailbud embryo"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             7 GlyValTyrPheAspGluAspPheHisValArgIleLeuAspValAspLysTyrAsnAla 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SerLysSerLeuGlnAspAsnThrAsnValPheIleAsnAsnIleGlnAsnMetGlnGly 46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    67 LeuLysAlaIleGlyLeuArgAsnArgValAlaAlaLeuSerGluGluArgLysArgLys 86
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40
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1 (bases 1 to 742)
Satou, Y., Shin-i, P., Kohara, Y. and Satoh, N.
Expressed genes in Ciona intestinalis (2002c)
Unpublished
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length:
Matches:
Conservative:
Mismatches:
Indels:
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                                                                                                                                                                                                                                                                       /organism="Ciona intestinalis"
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/db_xref="taxon:7719"
/clone="rcltb078f08"
                                                                                     Contact: Nori Satoh
Department of Zoology
Kyoto University
Sakyo-ku, Kyoto, Kyoto 606-8502, Japan
Tel: 81-75-753-4081
Fax: 81-75-705-1113
                                                                                                                                                                                                                                                                                                                                                                                                                                    265 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Xenopus laevis (African clawed frog)
Xenopus laevis
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                                                                                                                                                                                                                                    Location/Qualifiers
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/sex-"female"
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32.79%
29.50%
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919 541-4571
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                                     Xenopus laevis
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AW645594.1 GI:7403048
                                                              Kitayama, A., Terasaka, C., Mochii, M., Ueno, N., Shin-i, T. and Kohara
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GACGCGGGGCTGCACTTTGATGAGCTGAACAAATTGCGTATCTTGGACCCCGACGTCTCC 184
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CAGCAAACAACCGAATTAAAGGAGGAGTGCCGAGATTTTGTAGACAAAATAGGACATTTT 244
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245 CAGAAAGTAGTGGGGTGGACTAATTGAGCTTGTTGATGAGTTAGCTAAAGAAACGGAAAAC 304
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           85 ArglysGlnLysGluGlnGluArgMetLeuAlaGluLysGlnGluGluLeuGluArgLeu 104
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea; Pipidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25 AsnAlaSerLysSerLeuGlnAspAsnThrAsnValPheIleAsnAsnIleGlnAsnMet 44
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                                                                                                                                                                                                                                                                                                                                                                       /clone_lib="NIBB Mochii normalized Xenopus tailbud
library" 106 c 134 g 118 t
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                                                                                                                           Contact: Tadasu Shin-i
Center For Genetic Resource Information
National Institute of Genetics
1111 Yata, Mishima, Shizuoka 411-8540, Japan
Tel: 81-559-81-6856
Fax: 81-559-81-6855
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length:
Matches:
Conservative:
Mismatches:
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/organism="xenopus laevis"
/mol_type="mRNA"
/db_xref="taxon:8355"
/clone="xL078b07"
/tissue_type="whole embryo"
/dev_stage="stage 25"
                                                                                               Expressed genes in X. laevis embryo
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                                                                                                                                                                                                                            Email: tshini@genes.nig.ac.jp.
Location/Qualifiers
                                Kenopodinae; Xenopus.
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61.48%
32.79%
29.50%
                                               (bases 1 to 548)
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                                                                                                                Unpublished
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                                                                                                                                                                                                       Xenopodinae; Xenopus.

1 (Bases I to 609)

1 (bases I to 609)

Jr. Moore,D.T., Lai,W.S., Thorn,J.M., Kennington,E.A., Staffa,N.G. Jr., Moore,D.T., Bouffard,G.G., Beckstrom-Sternberg,S.M., Touchman,J.W., Bonaldo,M.F. and Soares,M.B.

The NIEMS Xenopus maternal EST project: interim analysis of the first 13,879 ESTs from unfertilized eggs

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Office of Clinical Research and Laboratory of Signal Transduction
National Institute of Environmental Health Sciences
A2-05 NIEHS, 101 Alexander Drive, Research Triangle Park, NC 27709,
                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Mesobatrachia; Pipoidea; Pipidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: black009@niehs.nih.gov
Clone is available through Research Genetics, Inc., 2130 Memorial
Parkway, Huntsville, AL 35901
phone 800-533-4363 ext.cdna, fax 256-536-9016 att:cdna, email
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA Sequencing and analyses performed by National Institutes of Health, Intramural Sequencing Center (NISC).
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Matches:
Conservative:
Mismatches:
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/cell_type="unfertilized egg"
/dev_stage="unfertilized egg"
Xenopus laevis (African clawed frog)
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Plate: 0153 row: F column:
Seq primer: T7 primer.
Location/Qualifiers
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83

us-09-866-582a-2.rst

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University of lowa 135 MEBRE, Iowa City, IA 52242, USA Tel: 319 335 8250
Fax: 319 335 8256
Email: bento-soares@ulowa.edu
The sequence contained an oligo-dT track that was present in the coligonucleotide that was used to prime the synthesis of first strand cDNA and therefore this may represent a bonafide poly A tail. CDNA Library Preparation: M.B. Soares Lab Clone distribution: clones will be available through Research Genetics (www.resgen.com) POLYA-Yes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EST 03-JUN-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              345 GAAGCTCAACAGCAGCAACTTCAAGCCCTAATAGCAGAAAAGAAAATGCAGCTAGAAAGG 404
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                                                                                                                                                                                                                                                                                                                                                                           44 MetGlnGlyLeuValAspLysTyrValSerAlaIleAspGlnGlnValGluArgLeuGlu 63
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bonaldo,M.F., Lennon,G. and Soares,M.B. Normalization and subtraction: two approaches to facilitate gene
                                                                                                                                                            4 ValAspArgGlyValTyrPheAspGluAspPheHisValArgIleLeuAspValAspLys
                                                                                                                                                                                                                                                                                                                     165 ACCCAGCAGACCATAGAGCTGAAGGAAGAGTGCAAAGACTTTGTGGACAAAATTGGCCAG
                                                                                                                                                                                                                                                                                                                                                                                                                               225 TITCAGAAAATAGTIGGIGGITIAAITGAGCIIGIIGAICAACIIGCAAAAGAAGCAGAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AI703692 427 bp mRNA linear EST 03-JUN-U1-R-AB1-yu-h-06-0-UI.s1 UI-R-AB1 Rattus norvegicus CDNA clone UI-R-AB1-yu-h-06-0-UI 3', mRNA sequence.
AI703692.1 GI:4991592
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     64 AlaGluLysLeuLysAlaIleGlyLeuArgAsnArgValAlaAlaLeuSerGluGluArg
                                                                                                                                                                                                                                                                     24 TyrAsnAlaSerLysSerLeuGlnAspAsnThrAsnValPheIleAsnAsnIleGlnAsn
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Coordinated Laboratory for Computational Genomics
0 0

    .427
    /organism="Rattus norvegicus"

Mismatches:
Indels:
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/strain="Sprague-Dawley"
                                                        Gaps:
                                                                                                      US-09-866-582A-2 (1-135) x BE884392 (1-917)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Rattus norvegicus (Norway rat)
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32.23%
29.50%
10
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                               Query Match:
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  Best Local
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/mol_type="mRNA"
/do_txref="taxon:9606"
/dolone="IMAGE:3907361"
/tissue_type="leiomyosarcoma"
/tissue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      вЕВВЯЗЭ2
601505670F1 NIH_MGC_71 Homo sapiens cDNA clone IMAGE:3907361 5',
mRNA sequence.
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                                                                                                                                                                                                                                                 CAGAAAGTAGTGGGTGGACTAATTGAGCTTGTTGATGAGTTAGCTAAAGAAACCGAAAAC 267
                                                                                                                                                                                                                                                                                                                                                                                                           85 ArgLysGlnLysGluGlnGluArgMetLeuAlaGluLysGlnGluGluLeuGluArgLeu 104
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NIH-WGC http://mgc.noi.nih.gov/.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CDNA Library Preparation: Life Technologies, Inc.
CDNA Library Preparation: Life Technologies, Inc.
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
Plate: LLAM9717 row: 1 column: 18
     CAGCAAACAACCGAATTAAAGGAGGAGTGCCGAGATTTTGTAGACAAAATAGGACATTTT
                                                                                                                                                                                           GlnGlyLeuValAspLysTyrValSerAlaIleAspGlnGlnValGluArgLeuGluAla
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Matches:
Conservative:
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Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 High quality sequence stop: 642.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unpublished
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Percent Similarity:
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BE884392
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// Clone_lib="Soares_mammary_gland_NMLMG"
// Clone_lib="Soares_mammary_gland_NMLMG"
// Clone_lib="Soares_mammary_gland by the modified polylinker; lst strand cDNA was prepared from mammary gland tissue from a lactating female, and was then primed with a Not I - oligo(dT) primer. Double-stranded cDNA was ligated to Eco RI adaptors (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pTTT3 vector. Library is normalized. Library was constructed by Bento Soares and M. Fatima Bonaldo. " a 122 c 74 g 139 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BY438168 RIKEN full-length enriched, 17 days embryo kidney Mus musculus cDNA clone I920155K14 3', mRNA sequence.
BY438168.1 GI:26721595
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                351 CAGCAGACTGTGGAGCTCAAGGAGGAGTGCAAGGACTTTGTGGACAAAATTGGCCAGTTT 292
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               Other_ESTs: ug64h07.yl
Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
MGI:952545.
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40
41
45
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Matches:
Conservative:
Mismatches:
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                                                                                                                                                                                                                                                                                 /sex="female (lactating)"
/tissue_type="mammary gland"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels:
                                                                                                                                                                                             /organism="Mus musculus"
/mol_type="mRNA"
/db_xref="taxon:10090"
                                                                                                                                                                                                                                                            /clone="IMAGE:1547197"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-866-582A-2 (1-135) x BE196900 (1-429)
                                                                                                                                                       Location/Qualifiers
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194.50
61.83%
30.53%
29.43%
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Best Local Similarity:
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VERSION
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BY438168
  JOURNAL
                                                                                                                                                     FEATURES
                        COMMENT
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         /clone="UI-R-ABI-yu-h-06-0-UI"
/dev_stage="adult"
/lab_host="DH10B (Life Technologies)"
/clone=lib="UI-R-ABI"
/note="Vector: p7730-Pac (Pharmacia) with a modified
/note="Vector: p7730-Pac (Pharmacia) with a modified
polylinker; Site_1: Not I; Site_2: Eco RI; The UI-R-ABI
library is a normalized library constructed from 16.5 dpc
rat ventricle. The tag is a string of 5 nucleotides
present between the Not I site and the oligo-dr track.
The library was constructed as described by Bonaldo,
Lennon and Soares, Genome Research 6: 791-806, 1996.
Tissue provided by Jim Lin, Department of Biology,
University of Iowa.
TAG_SED_None found"
111.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ug64h07.x1 Soares_mammary_gland_NNLMG Mus musculus cDNA clone IMAGE:1547197 3' similar to TR:Q61025 Q61025 HYPOTHETICAL 15.2 KD BR07EIN ; mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GlnMetGluGluGlnSerLeuIleLysValLysGlyGluGln-------Glu 119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        306 CAGAAAATCGTTGGTGGTCTGATTGAGCTTGTTGATCAGCTTGCCAAAGAAGCAGAGAAC 247
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ArgLysGlnLysGluGlnGluArgMetLeuAlaGluLysGlnGluLeuGluArgLeu 104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           366 CAGCAGACCACAGGAGCTCAAGGAAGAGGACCATTGTGGACAAAATTGGCCAGTTT 307
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5 AspArgGlyValTyrPheAspGluAspPheHisValArgIleLeuAspValAspLysTyr 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 (bases 1 to 429)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25 AsnAlaSerLysSerLeuGlnAspAsnThrAsnValPheIleAsnAsnIleGlnAsnMet
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Mismatches:
Indels:
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                                                                                                                                                                                                                                                                                                                                                                                                             Length:
Matches:
/db_xref="taxon:10116"
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Mus musculus
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BE196900/c
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ORIGIN
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TITLE
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KEYWORDS
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KEYWORDS SOURCE ORGANISM

REFERENCE AUTHORS

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/tissue_type="kidney"
/dev_stage="17 days embryo"
/clone_lib="RIKEN full-length enriched, 17 days embryo
                     /strain="C57BL/6J"
/db_xref="taxon:10090"
                                                                                           /clone="1920155K14"
/mol_type="mRNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AW533561.1 GI:7175975
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                                                                                                                                           Mammalis Eutherla; Kodentia; Sciurognathi; Muridae; Murinae; Musico; Mammalis; Eutherla; Kodentia; Sciurognathi; Muridae; Murinae; Mikaido,T., Osato,N., Saito,R., Suzuki,H., Yananaka!I., Kiyosawa,H., Yagi,K., Tomaru,Y., Hasegwa,Y., Nogami,A., Schonbach,C., Quacke,N., Schonbach,C., Gojobori,T., Baldarelli,R., Hill,D.P., Bult,C., Hume,D.A., Gojobori,T., Baldarelli,R., Hill,D.P., Bult,C., Hume,D.A., Batalov,S., Beisel,K.W., Blake,J.A., Bradat,D., Bursic,V., Chothia,C., Corbani,L.E., Cousins,S., Dalla,E., Dragani,T.A., Fletcher,C.F., Forrest,A., Frazer,K.S., Gaasterland,T., Gariboldi,M., Gissi,C., Godzik,A., Gough,J., Grimmond,S., Gustincich,S., Hirokawa,N., Jackson,I.J., Jarvis,E.D., Kanal,A., Kanal,H.H., Kawasawa,Y., Kedzierski,R.M., King,B.L., Konagaya,A., Kurochkin,I.V., Lee,Y., Lenhard,B., Lyons,P.A., Maltais,L., Marchionni,L., McKenzie,L., Miki,H.N., Nagashlma,T., Numate,K., Okido,T., Pavan,W.J., Pertea,G., Perrovsky,N., Pillai,R., Pontius,J.U., Qi,D., Ramachandran,S., Ravasi,T., Reed,J.C., Red,D.J., Reid,J., Ringwald,M., Sandelli,A., Schneider,C., Semple,C.A., Setou,M., Shimada,K., Sultana,R., Takenaka,Y., Taylor,M.S., Tasadale,R.D., Yang,I., Walls,C., Winshaw-Boris,A., Yang,I., Yang,I., Walls,C., Winsha,Y., Maki, M., Sakazume,N., Sato,K., Shirakl,T., Waki,K., Kawai,J., Azawa,K., Azawa,A., Yasunish,A., Xasunish,A., Xasunish,A., Xasunish,
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Normalization and subtraction of cap-trapper-selected cDNAs to prepare full-length cDNA libraries for rapid discovery of new genes. Genome Res. 10 (10), 1617-1630 (2000)
RIKEN Integrated sequence analysis (RISA) system--384-format sequencing pipeline with 384 multicapillary sequencer. Genome Res. 10 (11), 157-1771 (2000)
Computer-based methods for the mouse full-length cDNA encyclopedia: real-time sequence clustering for construction of a norredundant cDNA library. Genome Res. 11 (2), 281-289 (2011) cDNA library was prepared and sequenced in Mouse Genome Encyclopedia Project of Genome Exploration Research Group in Riken Genomic Sciences Center and Genome Science Laboratory in RIKEN. Division of Experimental Animal Research in Riken contributed to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: genome-res@gsc.riken.go.jp,
URL:http://genome.gsc.riken.go.jp,
Alzawa.K., Akimura,T., Arakawa,T., Carninci,P., Fukuda,S., Hirozane
Alzawa.K., Akimura,T., Arakawa,T., Carninci,P., Fonno,H., Miyazaki,A.,
T., Imotani,K., Ishli,Y., Itoh,M., Kawai,J., Konno,H., Sakai,K.,
Murata,M., Nakamura,M., Nomura,K., Numazaki,R., Ohno,M., Sakai,K.,
Sakazume,N., Sasaki,D., Sato,K., Shibata,K., Shiraki,T., Tagami
M., Waki,K., Watahiki,A., Muramatsu,M. and Hayashizaki,Y. Direct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Laboratory for Genome Exploration Research Group, RIKEN Genomic Sciences Center(GSC), Yokohama Institute The Institute of Physical and Chemical Research (RIKEN) 1-7-22 Suehiro-cho, Tsurumi-ku, Yokohama, Kanagawa 230-0045, Japan Fax: 81-45-503-9216
                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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Please visit our web site (http://genome.gsc.riken.go.jp) for
further details.
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                     Mus musculus (house mouse)
Mus musculus
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MEDLINE PUBMED

COMMENT

JOURNAL

TITLE

/organism="Mus musculus"

source

FEATURES

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AW533561 432 bp mRNA linear EST 06-MAR-2000 UI-R-BU0-and-d-03-0-UI.sl UI-R-BU0 Rattus norvegicus cDNA clone UI-R-BU0-and-d-03-0-UI 3', mRNA sequence.
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83 CAGCAGACTGTGGAGCTCCAAGGAGTGCAAGGACTTTGTGGACAAAATTGGCCAGTTT 142
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               105 GlnMetGluGluGlnSerLeuIleLysValLysGlyGluGln---------Glu 119
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Mammalia, Butheria, Rodentia, Sciurognathi, Muridae, Murinae,
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Bonaldo,M.F., Lennon,G. and Soares,M.B.
Normalization and subtraction: two approaches to facilitate gene
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University of Iowa
375 Newton Road , 4156 MEBRF, Iowa City, IA 52242, USA
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Mismatches:
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                                                                                                                                                                                                               Gaps:
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Rattus norvegicus
122 9
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EST 19-JUL-2001

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Coordinated Laboratory for Computational Genomics
University of Iowa
375 Newton Road, 4156 MEBRF, Iowa City, IA 52242, USA
375 Newton Road, 4156 MEBRF, Iowa City, IA 52242, USA
Tel: 319 335 9266
Email: bento-soares@uiowa.edu
The sequence contained an oligo-dT track that was present in the oligonucleotide that was used to prime the synthesis of first strand cDNA and therefore this may represent a bonafide polly A tail. The sequence tag present in the cDNA between the NotI site and the oligo-dT track served to identify it as a clone from the normalized ratk kidney pool library cDNA Library Preparation: M.B. Soares Lab Clone distribution: clones will be available through
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                for production of single-stranded circular bNA representing the pool of libraries. Single-stranded circular DNA representing these five normalized libraries was then used as a tracer in a subtractive hybridization with a driver (PCR amplified inserts from a plasmid DNA
                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
                                                                                                                                                                                                                                                                                                                                                    1 (bases 1 to 434)
Bonaldo,M.F., Lennon,G. and Soares,M.B.
Normalization and subtraction: two approaches to facilitate gene
                                                                                        BI289218
UI-R-DKO-cfj-c-07-0-UI.sl UI-R-DKO Rattus norvegicus cDNA clone
UI-R-DKO-cfj-c-07-0-UI 3', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /strain="Sprague-Dawley"
/db_xref="taxon:10116"
/clone="UT-R-DKO-cfj-c-07-0-UI"
/dev_stage="ADULT"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /organism="Rattus norvegicus"
66 TITATITICAGAAATGAACTGAAAATTTCAGT 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Genome Res. 6 (9), 791-806 (1996)
                                                                                                                                                                                                                                   Rattus norvegicus (Norway rat)
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                                                                                                                                                                                                                                                         Rattus norvegicus
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                                              RESULT 36
BI289218/c
                                                                                                              DEFINITION
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/db_xref="taxon:10]16"
/db_xref="taxon:10]16"
/db_bost="bullob dilte"
/dlone_lib="ul-R-Bu0"
/dlone_lib="ul-R-Bu0"
/dlone di ganglia dissues. For a detailed description of the
library from which this clone was derived, please visit
our web site at ratest.eng.ulowa.edu. The subtraction has
been previously described in (Bonaldo, Lennon and Soares,
Genome Research 6:791-806, 1996)
TAG_LIB=UT-R-Bu0
TAG_LIB=UT-R-Bu0
TAG_LIB=UT-R-Bu0
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                                       Email: bento-soares@ulowa.edu
The sequence contained an oligo-dT track that was present in the oligonucleotide that was used to prime the synthesis of first strand cDNA and therefore this may represent a bonafide poly A tail. The sequence tag present in the cDNA between the Noti site and the oligo-dT track served to identify it as a clone from the normalized ganglia library cDNA Library Preparation: M.B. Soares Lab Clone distribution: clones will be available through Research Genetics (www.resgen.com)
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                                                                                                                                                                                                                                                                                                                           1. 432
/organism="Rattus norvegicus"
/mol_type="mRRA"
/strain="Sprague-Dawley"
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Matches:
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114 c
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template preparation) comprising: a) a set of about 1,000 arrayed clones from each of the five non-normalized libraries of brain (CTOS), heart (CSOS), kidney (CUOS), aorta (CWOS), and placenta (CXOS). The resulting pool of approximately 5,000 clones represented about 33.3% of the final driver population. A set of about 2,000 arrayed clones from each of the five normalized libraries of brain (CTO), heart (CSO), kidney (CUO), aorta (CWO), and placenta (CXO). The resulting pool of about 10,000 clones represented about 66.8% of the final driver population. TAG_LIB-UI-R-DKO TAG_LISGED-TAT kidney pool 1745_SEOG-CAAGAGGGGTT TAG_LISGED-TAT Kidney pool 175 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             454 bp mRNA linear EST 21-JUN-2000 uf37b12.x1 Soares_mammary_gland_NNLMG Mus musculus cDNA clone IMAGE:1513535 3' Similar to TR:Q61025 Q61025 HYPOTHETICAL 15.2 KD PROTEIN. ;, mRNA sequence. BE132868 BE132868 GI:8595368 EST.
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Mus musculus
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Butheria; Rodentia; Sciurognathi; Muridae; Murinae; Musc. 1 (bases 1 to 454)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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Other_ESTs: uf37b12.y1
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Contact: Robert Strausberg, Ph.D.
Email: cgapbs-r@mail.nih.gov
This clone is available royalty-free through LLNL ; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
MGI:940387
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITATITITCAGAAATGAACTGAAAATTTCAGT 26
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Matches:
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                                                                                                                                                                                                                                                        High quality sequence stop: 414
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AI256584.1 GI:3864109
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194.50
61.83%
30.53%
29.43%
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AI170881 467 bp' mRNA linear EST 08-JAN-1999 EST216822 Normalized rat muscle, Bento Soares Rattus sp. cDNA clone RMUBC92 3' end, mRNA sequence.
AI170881.1 GI:4134559
                       ----Glu 119
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                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (pases 1 to 467)
Lee,N.H., Glodek,A., Chandra,I., Mason,T.M., Quackenbush,J.,
Kerlavage,A.R. and Adams,M.D.
Rat Genome Project: Generation of a Rat EST (REST) Catalog & Rat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /clone_lib="Normalized rat muscle, Bento Soares"
/note="Organ: muscle; Vector: pT7T3Pac; Site_l: EcoRI;
Site_2: NotI"
130 81 9 155 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             65 GluLysLeuLysAlaIleGlyLeuArgAsnArgValAlaAlaLeuSerGluGluArgLys
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gene Index
Unpublished
On Oct 6, 1998 this sequence version replaced gi:3710921.
Contact: Lee, NH
The Institute for Genomic Research
9712, Medical Center Drive, Rockville, MD 20850, USA
Tel: (301)-838-5229
Fax: (301)-838-0208
Email: nhlee@tigr.org
                                                                                                                                                                GlnMetGluGluGlnSerLeuIleLysValLysGlyGluGln----
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Mismatches:
Indels:
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Matches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /mol_type="mRNA"
/db_xref="taxon:10118"
/clone="RMUBC92"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
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Rattus sp.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /sex="female"
/dev_stage="adult"
/dev_stage="adult"
/lab_host="blub"
/clone_lib="Sugano mouse liver mlia"
/clone_lib="Sugano mouse liver mlia"
/clone_lib="Sugano mouse liver mlia"
/clone_lib="Site_lib="Site_lib="Site_lib="Site_lib="Site_lib="Site_lib="Site_lib="Site_lib="Site_lib="Site_lib="Site_lib="Site_lib="Site_lib="Site_lib="Site_lib="Site_lib="Site_lib="Site_lib="Site"
/ATGTGGCTTTTTTTTTTTTTTTTTTT]; double-stranded cDNA was
ligated to a Draili adaptor [TGTTGGCTACTGG], digested
and cloned into distinct Draili sites of the pME185-FL3
vector (5' site CACTGTGG, 3' site CACCATGGG, digested
and cloned into distinct Draili sites of the pME185-FL3
vector (5' site CACTGTGG, 3' site CACCATGGG, digested
and cloned into distinct DNA insert. Size selection was
performed to exclude fragments <1.5kb. Library
constructed by Dr. Sumio Sugano (University of Tokyo
Institute of Medical Science). Custom primers for
sequencing: 5' end primer CTGCTCTAAAAGCTGGG and 3' end
primer CGACCTGCAGCTCCAGCCAC."
97 a 127 c 76 g 163 t
                                                                                            Marra,M., Hillier,L., Allen,M., Bowles,M., Dietrich,N., Dubuque,T., Gelsel,S., Kucaba,T., Lacy,M., Le,M., Martin,J., Morris,M., Schellenberg,K., Steptoe,M., Tan,F., Underwood,K., Moore,B., Theising,B., Wylie,T., Lennon,G., Soares,B., Wilson,R. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            311 CAGAAAATFGTTGGTGGTCTAATTGAGCTTGTTGATCAGCTTGCCAAAGAAGAAGCAGAAC 252
                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
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                                                                                                                                                                                                                                                                                                                                                                                       Email: mouseest@watson.wustl.edu
This clone is available royalty.free through LLNL ; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
MGI:973804
                                                                                                                                                                                                                                           Unpublished
Contact: Marra M/Mouse EST Project
WashD-HHMI Mouse EST Project
WashD-HHMI University School of MedicineP
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108
Tel: 314 286 1800
Fax: 314 286 1810
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41
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Conservative:
Mismatches:
Indels:
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Location/Qualifiers
e 1. 463
/organism="Mus musculus"
/mol_type="mRNA"
/strain="C57BL"
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                                                                                                                                                                                                     Waterston, R.
The WashU-HHMI Mouse EST Project
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/clone="IMAGE:1889480"
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                 Mus musculus (house mouse)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          86 GCCCAGCAGCAGCTGCAGCCCTGATAGCAGAAAAGAAGAGCCGCAGCTAGAAAGGTAT 127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      105 GlnMetGluGluGlnSerLeuIleLysValLysGlyGluGln--------Glu 119
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           45 GlnGlyLeuValAspLysTyrValSerAlaIleAspGlnGlnValGluArgLeuGluAla 64
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  126 CGGGTTGAATATGAAGCTTTGTGTAAAGTAGAAGCAGAACAAAATGAATTTATTGACCAA 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Rodentia, Sciurognathi, Muridae, Murinae,
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The sequence contained an oligo-dT track that was present in the oligonucleotide that was used to prime the synthesis of first
oligo-dT priming.Size-selected by column fractionation; average insert size 1.06kb. Primary library, unamplified." 79 g 160 t 1 others
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Bonaldo,M.F., Lennon,G. and Soares,M.B.
Normalization and subtraction: two approaches to facilitate gene
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UI-R-DY1-cnu-j-06-0-UI.S1 UI-R-DY1 Rattus norvegicus cDNA clone
UI-R-DY1-cnu-j-06-0-UI 3', mRNA sequence.
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Conservative:
Mismatches:
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                                                                                                                                                                        Pred.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BM876762 474 bp mRNA linear EST 12-MAR-2002 477 bb mRNA linear EST 12-MAR-2002 4770c01.XI.Melton Mouse Els Fancheas Library 2 M1682 Mus musculus cDNA clone IMAGE:568164 3' similar to TR:061025 Q61025 HYPOTHETICAL 15.2 KD PROTEIN.; mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mus musculus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.

[ (bases 1 to 474)
Melton,D., Brown,J., Kenty,G., Permutt,A., Lee,C., Kaestner,K.,
Lemishka,I., Scearce,M., Brestell,J., Gradvohl,G., Clifton,S.,
Hillier,L., Marra,M., Pape,D., Wylie,T., Martin,J., Blistain,A.,
Schmitt,A., Theising,B., Ritter,E., Ronko,I., Bennett,J., Cardenas,
Jackson,Y. and Bowers,Y.
Endocrine Pancreas Consortium
Unpublished
                                                                                                                                                                                                                      GCCCAGCAACAGCAACTGCAGGCACTGATAGCAGAAAAGAAGATGCAGCTAGAAAGGTAT 110
                                                                                                                                                                                                                                                                                       GInMetGluGluGlnSerLeuIleLysValLysGlyGluGln------Glu 119
                                           GAGAAAATGAAGGCCATTGGTGCTCGAAACTTGCTGAAATCCATAGCGAAGCAGGAGAA 170
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Endocrine Pańcreas Consortium
Barvard University, Howard Hughes Medical Institute
Dept of Molecular and Cellular Biology, 7 Divinity Ave, Cambridge,
MA 02138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (brownefas.harvard.edu)
MGI:1959542 This sequence now available from the IMAGE consortium,
for clone orders contact: info@image.llnl.gov
                                                                                                                                                                                              ArgLysGlnLysGluGlnGluArgMetLeuAlaGluLysGlnGluGluLeuGluArgLeu 104
                                                                                                     84
            GlnGlyLeuValAspLysTyrValSerAlaIleAspGlnGlnValGluArgLeuGluAla 64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: dmelton@biohp.harvard.edu
Library was constructed by Dr. Douglas Melton DNA sequencing by:
Washington University Genome Sequencing Center For information o
Obtaining a clone please contact: Juliana Brown
                                                                                                     GluLysLeuLysAlaIleGlyLeuArgAsnArgValAlaAlaLeuSerGluGluArgLys
                                                                                                                                                                                                                                                                                                                                  CGGGTTGAATATGAAGCTTTGTGTAAAGTAGAAGCAGAACAAAATGAATTTATTGACCAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Contact: Douglas Melton, Klaus H. Kaestner, & Hiroshi Inoue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /tissue_type="Total pancreas"
/dev_stage="Embryonic day 16.5"
/lab_host="TOP10"
                                                                                                                                                                                                                                                                                                                                                                                  120 LeuMetIleGlnLysLeuSerAspSerSerSer 130
                                                                                                                                                                                                                                                                                                                                                                                                               TTTATTTTCAGAAATGAACTGAAAATTTCAGT 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /organism="Mus musculus"
/mol_type="mRNA"
/strain="ICR"
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High quality sequence stop: 466.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /sex="Both"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 rel: 617-495-1812
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Fax: 617-495-8557
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BM876762/c
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KEYWORDS
SOURCE
ORGANISM
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JOURNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REFERENCE
AUTHORS
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Page 25

us-09-866-582a-2.rst

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BQ174785 539 bp mRNA linear EST 30-APR-2002 UI-M-DJ2-bwb-d-13-0-UI.s1 NIH_BMAP_DJ2 Mus musculus cDNA clone UI-M-DJ2-bwb-d-13-0-UI 3', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. I (bases 1 to 539)
Bonaldo, M. F., Lennon, G. and Soares, M. B.
Normalization and subtraction: two approaches to facilitate gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tissue Procurement: Dr. Robin Davisson cones, University of Iowa CDNA Library preparation: Dr. M. Bento Soares, University of Iowa CDNA Library Arrayed by: Dr. M. Bento Soares, University of Iowa Clone Distribution: Researchers may obtain clones from Research Genetics (www.resgen.com).
      127 CGGGTTGAATATGAAGCTTTGTGTAAAGTAGAAGCAGAACAAAATGAATTTATTGACCAA 68
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Chin, H
National Institute of Mental Health
National Institute of Mental Health
2001 Executive Blvd. Room 7N-7190, MSC 9643, Bethesda, MD
20892-9643, USA
Tel: 301 443 1706
Fax: 301 443 9890
Email: mEST@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TAG_TISSUE=subfornical organ and postrema
TAG_SEQ=GCTACATGAT"
144 c 103 g 177 t
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                                                  120 LeuMetIleGlnLysLeuSerAspSerSerSer 130
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Matches:
                                                                           /organism="Mus musculus"
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1. .539
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Mus musculus
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BQ174785.1 GI:20350277
EST.
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194.50
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Pred. No.:
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                                                                                                                                                           BQ174785/c
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TITLE
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PUBMED
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         g
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                                                                                                                                                                                                                                  /organism="Rattus norvegicus"
/mol_type="mrNA"
/strain="Sprague-Dawley"
/db_xref="taxon:1016"
/clone="UI-R-DYI-cnu-j-06-0-UI"
/tlssue_type="Cartilage"
/dev_stage="ADULT"
/tlssue_type="Cartilage"
/dev_stage="ADULT"
/tlssue_type="Cartilage"
/clone_lib="UI-R-DYI"
/rote="Organ: Femur and Tibia; Vector: pT7T3D-Pac
/clone_lib="UI-R-DYI"
/rote="Corgan: Femur and Tibia; Vector: pT7T3D-Pac
/pharmacia) with a modified polylinker; Site=1: Not I;
Site_2: Eco RI: UI-R-DYI is a normalized CDNA library
containing the following tissue(s): Rat Cartilage from
Femur and Tibia. The library was constructed according to
Bonaldo, Lennon and Soares, Genome Research, 6:791-806,
1996. First strand CDNA synthesis was primed with an
oligo-dT primer containing a Not I site. Double stranded
cDNA was ligated to an Eco R I adaptor: dispested with Not
I, and cloned directionally into pT7T3-Pac vector. The
oligonucleotide used to prime the synthesis of
first-strand cDNA contains a library tag sequence that is
located between the Not I site and the (dT)18 tail. The
sequence tag for this library is CTAATGGACG. The Rat
cartillage tissue was provided by Dr Jeff Stevens at the
University of Towa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ::: ||||
367 CAGCAGACCACAGAGGTGCAAGGACTTTGTGGGAAATTGGCCAGTTT 308
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    307 CAGAAAATCGTTGGTGGTCTGATTGAGCTTGTTGATCAGCTTGCCAAAGAAGCAGGAGAAC 248
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             247 GAGAAAATGAAGGCCATTGGTGCTCGAAACTTGCTGAAATCCATAGCGAAGCAGAGAA 188
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strand cDNA and therefore this may represent a bonafide poly A tail. The sequence tag present in the cDNA between the NotI site and the oligo-dT track served to verify it as a clone from the normalized cartilage library ONA Library Preparation: M.B. Soares Lab Clone distribution: clones will be available through Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25 AsnAlaSerLysSerLeuGlnAspAsnThrAsnValPheIleAsnAsnIleGlnAsnMet 44
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                64
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41
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Matches:
Conservative:

    .517
    /organism="Rattus norvegicus"

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TAG_LIB=UI-R-DYI
TAG_TISSUE-cartilage
TAG_SEQ-CTAATGGACG"
                                                                                                                                                                                Location/Qualifiers
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                                                                                                             Genetics (www.resgen.com)
Seg primer: M13 Forward
POLYA-Yes.
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194.50
61.83%
30.53%
29.43%
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Mus musculus

Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Eutheria; Rodentia; Sclurognathi; Muridae; Murinae; Mus.

1 (Dases 1 to 573)

Tanaka,T.S., Jaradat,S.A., Lim,M.K., Kargul,G.J., Wang,X., Grahovac,M.J., Pantano,S., Sano,Y., Piao,Y., Nagaraja,R., Doi,H., Wood,W.H.

Genome-wide expression profiling of mid-gestation placenta and embryo using a 15,000 mouse developmental cDNA microarray

Proc. Natl. Acad. Sci. U.S.A. 97 (16), 9127-9132 (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AW553522 ST3 bp mRNA linear EST 10-JUN-2003 L0228G10-3 NIA Mouse Newborn Ovary cDNA Library Mus musculus cDNA AW553522 GI:31568866
                                                                                                                                                         307 CAGAAAATTGTTGGTGGTCTAATTGAGCTTGTTGATCAGCTTGCCAAAGAAGAAGCAGAAAC 248
                                                                                                                                                                                                                                                                                                                                                                                    ArgLysGlnLysGluGlnGluArgMetLeuAlaGluLysGlnGluGluLeuGluArgLeu 104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 105 GlnMetGluGluGlnSerLeuIleLysValLysGlyGluGln-------Glu 119
                                                                                                                                   5 AspArgGlyValTyrPheAspGluAspPheHisValArglleLeuAspValAspLysTyr 24
                                                                                                                                                                                                                                                                                                64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25 AsnalaSerLysSerLeuGlnAspAsnThrAsnValPhelleAsnAsnIleGlnAsnMet
                                                                                                                                                                                                                                                CAGCAGACTGTGGAGCTCAAGGAGGAGTGCAAGGACTTTGTGGACAAAATTGGCCAGTTT
                                                                                                                                                                                                                                                                                      GlnGlyLeuValAspLysTyrValSerAlaIleAspGlnGlnValGluArgLeuGluAla
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                National Institute on Aging/National Institutes of Health 333 Cassell Drive, Suite 4000, Baltimore, MD 21224-6820, USA Email: cdnaelgsun.grc.nia.nih.gov
Plate: L0228 row: C. column: 10
Seq primer: -21M3 Forward
High quality sequence stop: 573
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      10922068
On Mar 7, 2000 this sequence version replaced gi:7198945
       Conservative:
Mismatches:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          120 LeuMetIleGlnLysLeuSerAspSerSerSer 130
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/mol_type="mRNA"
/strain="C57BL/G1"
/db_xref="niaEST:L0228C10-3"
/db_xref="taxon:10090"
                                         Indels:
                                                                                               US-09-866-582A-2 (1-135) x BQ174785 (1-539)
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Laboratory of Genetics
   61.83%
30.53%
29.43%
Percent Similarity:
Best Local Similarity:
Query Match:
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AGENCOURT_10617261 NIH_MGC_169 mRNA linear EST 21-OCT-2002 IMAGE:6742597 5', mRNA sequence.
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440
41
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Matches:
Conservative:
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/dev_stage="Newborn Ovary"
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194.50
61.83%
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Query Match:
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27

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/clone="Indacolity".
/lab_host="Indacolity".
/lab_host="DH108 (phage-resistant)"
/clone=lib="NHH_MGC_164"
/clone=lib="NHH_MGC_164"
/note="Vector: pCMV-SPORT6"
NotI; Non-normalized full-length enriched library from pooled mouse embryonic limb, maxilla and mandible, day 10.5 and 11.5 (size selected for the 0.5-1 kb fragments) cloned directionally, priming method: Oligo-dT. cDNA enrichment: >lk bp, Average insert size 1.8k bp. Priming sequence: 5 GACTAGTTCTAGATCGCGAGGCGCCCC(T) 3'. Tissue contributed by, David Rowe: Library constructed by ResGen, Invitrogen Corp."
Invitrogen Corp."
                                                                                                                                                                                                                                                                               EST 06-JAN-2003
   265 CAGCAGACTGTGGAGCTCAAGGAGGAGTGCAAGGACTTTGTGGACAAAATTGGCCAGTTT 324
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus I (bases 1 to 633)
NIH-WGC http://mgc.ncl.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CDNA Library Preparation: Invitrogen Corporation Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Agencourt Bloscience Corporation Clone distribution: MGC clone distribution information can be found through the I.M.A.G.E. Consortium/LLNL at: http://mage.llnl.gov
Plate: NDAMOGG row: b column: 02
High quality sequence stop: 633.
Location/Qualifiers
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                                                                                                                                                                                                                                                                      CA977210
AGENCOURT_11292396 NIH_MGC_164 Mus musculus cDNA clone IMAGE:30147961 5', mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Robert Strausberg, Ph.D.
Email: cgapbs-rémail.nih.gov
Tissue Procurement: Dr. David Rowe and Dr. Mina
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Matches:
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Mismatches:
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/db_xref="taxon:10090"
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VERSION
KEYWORDS
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AUTHORS
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                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. I thases 1 to 615)
NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished
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Email: Ggapbs-remail.nih.gov
Tissue Procurement: Dr. Jonathan Kuo, NIMH
CDNA Library Preparation: Michael Brownstein Laboratory
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Agencourt Biosocience Corporation
Clone distribution: MGC clone distribution information can be
found through the I.M.A.G.E. Consortlum/LLNL at:
http://image.lnl.gov
Plate: LLCM3080 row: o column: 12
High quality sequence stop: 540.
Location/Qualifiers

cee /organism="Mus musculus"
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Mismatches:
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Matches:
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/clone="IMAGE:6742597"
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                                                     Mus musculus (house mouse)
Mus musculus
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   BU961938:1 GI:24191510
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TAG_TISSUE=duodenum
TAG_SEQ=TGTGGTTCAT"
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                                                                                                                                                                                    85 ArgLysGlnLysGluGlnGluArgMetLeuAlaGluLysGlnGluGluLeuGluArgLeu 104
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The sequence contained an oligo-dT track that was present in the oligonucleotide that was used to prime the synthesis of first strand cDNA and therefore this may represent a bonafide poly A tail. The sequence tag present in the cDNA between the NotI site and the oligo-dT track served to identify it as a clone from the normalized duodenum library cDNA Library Preparation: M.B. Soares Lab Clone distribution: clones will be available through Research Genetics (www.resgen.com)

Seq primer: M13 Forward
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Bonaldo,M.F., Lennon,G. and Soares,M.B.
Normalization and subtraction: two approaches to facilitate gene
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UI-R-EPO-COG-b-22-0-UI.S1 UI-R-EPO Rattus norvegicus CDNA clone
BQ209888
                                                                                                                                                                                                                         445 GCCCAGCAGCAGCCAGGCCCTGATAGCAGAAAAGAAGAAGGCAGCTAGAAAGGTAT
                                                                             GluLysLeuLysAlaIleGlyLeuArgAsnArgValAlaAlaLeuSerGluGluArgLys
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
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375 Newton Road , 4156 MEBRF, Iowa City, IA 52242,
17E1: 319 335 9565
Fax: 319 335 9565
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    /organism="Rattus norvegicus"

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97044477
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Rattus norvegicus
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Coordinated Laboratory
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H3085G04-3 NIA Mouse 15K cDNA Clone Set Mus musculus cDNA clone H3085G04 3', mRNA sequence.
BG070290
BG070290.1 GI:12552859
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Genome-wide expression profiling of mid-gestation placenta and embryo using a 15,000 mouse developmental cDNA microarray Proc. Natl. Acad. Sci. U.S.A. 97 (16), 9127-9132 (2000)
directionally into pT7T3-Pac vector. The oligonucleotide used to prime the synthesis of first-strand cDNA contains a library tag sequence that is located between the Not I site and the (dT)18 tail. The sequence tags for this library are: distal colon, GAAGTGCTCC; osteoblast, AAGATATCAA; cell line R3327 5A, GGACTAGATC; cell line R3327 5P, CACGTGAGAT; duodenum, TGTGGTTCAT; prostate,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        127 GAGGCAGGGCTGCACTTCGATGAGCTGAACAAGCTTCGGGTGTTGGACCCAGAGGTTACC 368
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      367 CAGCAGACCACAGAGCTCAAGGAAGAGTGCAAGGACTTTGTGGGAAAAATTGGCCAGTTT 308
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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/organism="Mus musculus"
/mol_type="mrNA"
/strain="cSTBL/60"
/db_xref="taxon:10090"
/db_xref="taxon:10090"
/clone="UI-W-CGOp-biq-b-11-0-UI"
/lab_host="UH-BMAP_Ret4_S2"
/rlote="vector: pT7130-Pac (Pharmacia) with a modified
polylinker; Site_1: Not I; Site_2: Eco RI; The
NIH_BMAP_Ret4_S2 library is a subtracted library,
ultimately derived from mouse retina tissue libraries at
various stages of development. For a detailed description
of the library from which this clone was derived, please
visit our web site at brainest.eng.uiowa.edu. The tissue
for this library was contributed by Dr. Xin-Yuan Fu, Yale
University School of Medicine"
185 a 185 c 148 g 199 t 1 others
                                                                                                                                                                                                                                                                                                               UI-M-CGOp-biq-b-11-0-UI.rl NIH_BMAP_Ret4_S2 Mus musculus cDNA clone UI-M-CGOp-biq-b-11-0-UI.5', mRNA sequence.
                                                                                                                     CDNA Library preparation: Dr. M. Bento Soares, Univeristy of Iowa CDNA Library Arrayed by: Dr. M. Bento Soares, Univeristy of Iowa DNA Sequencing by: Dr. M. Bento Soares, Univeristy of Iowa Clone Distribution: Researchers may obtain clones from Research Genetics (www.resgen.com).

Seq primer: M13 REVERSE.

Location/Qualifiers
                            GlnMetGluGluGlnSerLeuIleLysValLysGlyGluGln-------Glu 119
    85 ArgLysGlnLysGluGlnGluArgMetLeuAlaGluLysGlnGluGluLeuGluArgLeu 104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. I (bases 1 to 691)
Bonaldo, M.F., Lennon, G. and Soares, M.B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Chin, H
National Institute of Mental Health
6001 Executive Blvd. Room 7N-7190, MSC 9643, Bethesda, MD
20892-9643, USA
Tel: 301 443 9890
Fax: 301 443 9890
Fmail: mEsT@mail.nih.gov
Tissue Procurement: Dr. Xin-Yuan Fu, Yale University School of Medicine
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Conservative:
Mismatches:
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Genome Res. 6 (9), 791-806 (1996)
97044477
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Mus musculus
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BM940927.1 GI:19400079
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/clone_libst__NIA Mouse 15K cDNA Clone Set"
/note="Vector: pSPORT1; Site_1: Sal1; Site_2: NotI; This clone is among a rearrayed set of 15,247 clones from 11 embryo cDNA libraries (including preimplantation stage embryos from unfertilized egg to blastocyst, embryonic part of ET.5 embryos, extraembryonic part of ET.5 embryos, and El2.5 female mesonephros/gonad) and one newborn ovary cDNA library. Average insert size 1.5 kb. All source libraries are cloned unidirectionally with Oligo(dT) Not primers. References include: (1) Genome-wide expression profiling of mid-gestation placenta and embryo using a 15,000 mouse developmental cDNA microarray, 2000, Proc. Natl. Acad. Sci. U S A, 97: 9127-9128; (2)
Large-scale cDNA analysis reveals phased gene expression patterns during preimplantation mouse development, 2000, Development, 127: 1737-1749; (3) Genome-wide mapping of unselected transcripts from extraembryonic tissue of 7.5-day mouse embryos reveals enrichment in the t-complex and under-representation on the X chromosome, 1998, Hum Mol Genet 7: 1967-1978.
                                                                                                                                         Please
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    45 GlnGlyLeuValAspLysTyrValSerAlaIleAspGlnGlnValGluArgLeuGluAla 64
                                                                                                                                                                                                                                                                                                                                                                                                                                                             /sex="Clones arrayed from a variety of cDNA libraries"
/dev_stage="Clones arrayed from a variety of cDNA
libraries"
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                Contact: George J. Kargul
Laboratory of Genetics
National Institute on Aging/National Institutes of Health
333 Cassell Drive, Suite 4000, Baltimore, MD 21224-6820, USA
Email: cdna@lgsun.grc.nia.nih.gov
This clone set has been freely distributed to the community. For the First here: All Solve Seq primer: -21M13 Forward
High quality sequence stop: 687
POLYA-Yes.
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Matches:
Conservative:
Mismatches:
Indels:
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                                                                                                                                                                                                                                                                                                   1. .687
/organism="Mus musculus"
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174 c 136 g 2
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/clone="H3085G04"
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                                                                                                                                                                                                                                                                           Location/Qualifiers
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/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /lab_host="DH10B"
  Other_ESTs: H3085G04-5
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Best Local Similarity:
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COMMENT
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BQ177222/c DEFINITION

RESULT 49

ACCESSION VERSION KEYWORDS SOURCE

ORGANISM

REFERENCE AUTHORS TITLE

MEDLINE PUBMED COMMENT

FEATURES

JOURNAL

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/de_stage="Adult"
//de_stage="Adult"
//de_stage="Adult"
//de_stage="Adult"
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//doc=="Organ: brain; Vector: pT7T3-Pac (Pharmacia) with a modified polylinker; Site_1: EcoR I; Site_2: Not I; U1-M-DJ2 is a subtracted cDNA library constructed according to Bonaldo, Lennon and Soares, Genome Research, 6:791-806, 1996. First strand cDNA synthesis was primed with an oligo-off primer containing a Not I site. Double stranded cDNA was ligated to an EcoR I adaptor, digested with Not I, and cloned directionally into pT7T3-Pac vector. The oligonucleotide used to prime the synthesis of first-strand cDNA contains a library tag sequence that is located between the Not I site and the (dT)18 tail. The sequence tag for this library is: GCTACATGAT, subfornical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              H3136B04-3 NIA Mouse 15K cDNA Clone Set Mus musculus cDNA clone H3136B04 3', mRNA sequence.
BG074519 GI:12557088
EST.
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Mus musculus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            45 GlnGlyLeuValAspLysTyrValSerAlaIleAspGlnGlnValGluArgLeuGluAla 64
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307 CAGAAAATIGTIGGIGGICIAAFIGAGCTIGTIGATCAGCTIGCCAAAGAAGCAGAGAAC
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/tissue_type="subfornical organ and postrema"
                                                                                                                                                                                                                                                                                                                                                                                         organ and area postrema.
TAG_LIB=UI-M-DJ2
TAG_TISSUB=subfornical organ and postrema
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Best Local Similarity:
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BG074519/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Email: mEST@mail.nih.gov
Tissue Procurement: Dr. Robin Davisson
CDNA Library preparation: Dr. M. Bento Soares, University of Iowa
CDNA Library Arrayed by: Dr. M. Bento Soares, University of Iowa
DNA Sequencing by: Dr. M. Bento Soares, University of Iowa
Clone Distribution: Researchers may obtain clones from Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GlnMetGluGluGluGlnSerLeuIleLysValLysGlyGluGln------Glu 119
                                                                                                                                                                                                                                                                                                                           232 GAGAAGATGAAGGCCATTGGTGCTCGGAACTTGCTGAAGTCCATAGCGAAGCAGAGAAA 173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           352 CAGCAGACTGTGGAGCTCAAGGAGGAGTGCAAGGACTTTGTGGACAAAATTGGCCAGTTT 293
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                             25 AsnAlaSerLysSerLeuGlnAspAsnThrAsnValPheIleAsnAsnIleGlnAsnMet 44
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                                                                                                 5 AspargGlyValTyrPheAspGluAspPheHisValArgIleLeuAspValAspLysTyr 24
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                                                                                                                                                                                                                                                                                            45 GlnGlyLeuValAspLysTyrValSerAlaIleAspGlnGlnValGluArgLeuGluAla
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="Mus musculus"
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Contact: Chin, H
National Institute of Mental Health
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Genome Res. 6 (9), 791-806 (1996)
       Gaps:
                                                   US-09-866-582A-2 (1-135) x BM940927 (1-691)
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Seg primer: M13 FORWARD
POLYA=Yes.
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Mus musculus
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/clone_lib=_nia Mouse 15k cDNA Clone Set"
/clone_lib=_nia Mouse 15k cDNA Clone Set"
/clone is among a rearrayed set of 15,247 clones from 11
clone is among a rearrayed set of 15,247 clones from 11
embryo cDNA libraries (including preimplantation stage embryos from unfertilized egg to blastocyst, embryonic part of E7.5 embryos.

, and E12.5 female mesonephros/gonad) and one newborn ovary cDNA library. Average insert size 1.5 kb. All source libraries are cloned unidirectionally with Oligo(dT) Not primers. References include: (1) Genome-wide expression profiling of mid-gestation placenta and embryo using a 15,000 mouse developmental cDNA microarray, 2000, Proc. Natl. Acad. Sci. U S A, 97: 9127-9132; (2)
Large-scale cDNA analysis reveals phased gene expression partering alternation place.
                                                                                                                                                                                                                                                                             Contact: George J. Kargul
Laboratory of Genetics
National Institute on Aging/National Institutes of Health
National Institute on Aging/National Institutes of Health
Email: acasell Drive, Suite 4000, Baltimore, MD 21224-6820, USA
Email: cdna@lgsun.grc.nia.nih.gov
This clone set has been freely distributed to the community. Please
Plate: H3136 row: B column: 04
Seq primer: -21M13 Forward
High quality sequence stop: 775
POLYA-Yes.
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              Tanaka T.S., Jaradat, S.A., Lim,M.K., Kargul,G.J., Wang,X., Grahovac, M.J., Pantano,S., Sano,Y., Piao,Y., Nagaraja,R., Doi,H., Wood,W.H.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /sex="Clones arrayed from a variety of cDNA libraries"
dev_stage="Clones arrayed from a variety of cDNA
                                                                                                       Genome-wide expression profiling of mid-gestation placenta and embryo using a 15,000 mouse developmental cDNA microarray Proc Natl. Acad. Sci. U.S.A. 97 (16), 9127-9132 (2000) 20381348
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Matches:
Conservative:
Mismatches:
Indels:
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/organism="Mus musculus"
/mol_type="mRNA"
/strain="c57BL/6"
/db_xref="niaEST:H3136B04-3"
/db_xref="naxon:10090"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /clone="H3136B04"
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                                                                                                                                                                                                                                                         Other_ESTs: H3136B04-5
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289 CAGAAAATTGTTGGTGGTCTAATTGAGCTTGTTGATCAGCTTGCCAAAGAAGCAGAGAAC 230
                                                                                                                 229 GAGAAGATGAAGGCCCATTGGTGCTCGGAACTTGCTGAAGTCCATAGCGAAGCAGAGAGAA 170
                                                                                                                                                                             105 GlnMetGluGluGlnSerLeuIleLysValLysGlyGluGln.........Glu 119
                                                                                                                                                        85 ArgLysGlnLysGluGlnGluArgMetLeuAlaGluLysGlnGluGluLeuGluArgLeu 104
                                                                                                                                                                                                                                                            64
GlnGlyLeuValAspLysTyrValSerAlaIleAspGlnGlnValGluArgLeuGluAla
                                                                            65 GluLysLeuLysAlaIleGlyLeuArgAsnArgValAlaAlaLeuSerGluGluArgLys
                                                                                                                                                                                                                                                                                                                   120 LeuMetIleGlnLysLeuSerAspSerSerSer 130
                                                                                                                                                                                                                                                                                                                                              49 TTTATTTTCAGAAATGAACTGAAAATTTCAGT 17
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